1011

OIPE

#_3

RAW SEQUENCE LISTING DATE: 10/04/2001 PATENT APPLICATION: US/09/863,063 TIME: 17:42:45

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3 <110> APPLICANT: GREENSTEIN, DAVID

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MILLER, MICHAEL A.
      6 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS OF NEMATODE CONTROL
      8 <130> FILE REFERENCE: N-7088
     10 <140> CURRENT APPLICATION NUMBER: 09/863,063
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C--> 11 <141> CURRENT FILING DATE: 2001-09-21
     13 <150> PRIOR APPLICATION NUMBER: 60/205,829
     14 <151> PRIOR FILING DATE: 2000-05-19
     16 <150> PRIOR APPLICATION NUMBER: 60/274,358
     17 <151> PRIOR FILING DATE: 2001-03-08
     19 <160> NUMBER OF SEQ ID NOS: 33
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                                          25
     35 Val Ile Asn Ser Ser Ala Arg Arg Ile Gly Tyr Gly Ile Lys Thr Thr
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                                     40
     38 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Leu Asp Pro
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     41 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly
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     44 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Val Glu Trp Thr Asn Thr
     47 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
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     63 Ile Val Phe Asn Ala Pro Tyr Asp Asp Lys His Thr Tyr His Ile Lys
                                         25
                     20
     66 Val Ile Asn Ser Ser Ala Arg Arg Ile Gly Tyr Gly Ile Lys Thr Thr
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     67
     69 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Leu Asp Pro
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72 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly

Input Set : A:\N-7088us.app

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75 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Val Glu Trp Thr Asn Thr
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78 Pro Asp Gly Ala Ala Arg Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
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81 Gly Met Val Arg Arg Lys Asn Leu Pro Ile Glu Tyr Asn Pro
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94 Ile Val Phe Asn Ala Pro Tyr Asp Asp Lys His Thr Tyr His Ile Lys
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97 Val Ile Asn Ser Ser Ala Arg Arg Ile Gly Tyr Gly Ile Lys Thr Thr
98
            35
100 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Leu Asp Pro
101
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                             55
103 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly
                         70
                                              75
104 65
106 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Val Glu Trp Thr Asn Thr
107
                     85
109 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
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128 Val Ile Asn Ser Ser Ala Arg Arg Ile Gly Tyr Gly Ile Lys Thr Thr
131 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Leu Asp Pro
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134 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly
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137 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Val Glu Trp Thr Asn Thr
138
140 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
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156 Ile Val Phe Asn Ala Pro Tyr Asp Asp Lys His Thr Asp His Ile Lys
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157
                 20
159 Val Ile Asn Ser Ser Ala Arg Arg Ile Gly Tyr Gly Ile Lys Thr Thr
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162 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Phe Asp Pro
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163
165 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly
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                                              75
166 65
168 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Val Glu Trp Thr Asn Thr
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169
171 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
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187 Ile Val Phe Asn Ala Pro Tyr Asp Asp Lys His Thr Tyr His Ile Lys
                                      25
                 20
190 Val Ile Asn Ser Ser Ala Arg Arg Ile Gly Tyr Gly Ile Lys Thr Ile
             35
191
193 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Leu Asp Pro
194
         50
196 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly
                                              75
                         70
197 65
199 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Val Glu Trp Thr Asn Thr
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202 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
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203
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205 Gly Met Val Arg Arg Lys Asn Leu Pro Ile Glu Tyr Asn Pro
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209 <210> SEQ ID NO: 7
210 <211> LENGTH: 126
211 <212> TYPE: PRT
212 <213> ORGANISM: Caenorhabditis elegans
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218 Ile Val Phe Asn Ala Pro Tyr Asp Asp Lys His Thr Tyr His Ile Lys
                                                          30
219
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221 Val Ile Asn Ser Ser Ala Arg Arg Ile Val Tyr Gly Ile Lys Thr Thr
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224 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Leu Asp Pro
225
         50
                             55
227 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly
228 65
                                              75
                         70
230 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Val Glu Trp Thr Asn Thr
231
                     85
233 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
234
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236 Gly Met Val Arg Arg Lys Asn Leu Pro Ile Glu Tyr Asn Pro
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242 <212> TYPE: PRT
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      1
249 Ile Val Phe Asn Ala Pro Tyr Asp Asp Lys His Thr Tyr Arg Ile Lys
252 Val Ile Asn Ser Ser Ala Arg Arg Ile Gly Tyr Gly Ile Lys Thr Thr
253
             35
255 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Leu Asp Pro
256
         50
                             55
258 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly
                         70
                                              75
261 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Val Glu Trp Thr Asn Thr
262
                     85
264 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
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265
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267 Gly Met Val Arg Arg Lys Asn Leu Pro Ile Glu Tyr Asn Pro
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271 <210> SEQ ID NO: 9
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280 Ile Val Phe Asn Ala Pro Tyr Asp Asp Lys His Thr Tyr His Ile Lys
281
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                                      25
283 Val Ile Asn Ser Ser Ala Arg Arg Ile Gly Tyr Gly Ile Lys Thr Thr
284
             35
286 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Leu Asp Pro
287
         50
                             55
289 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly
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Input Set : A:\N-7088us.app

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75
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                         70
290 65
292 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Val Glu Trp Thr Asn Thr
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293
295 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
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298 Gly Met Ala Arg Arg Lys Asn Leu Pro Ile Glu Tyr Asn Pro
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311 Ile Val Phe Asn Ala Pro Tyr Asp Asp Lys His Thr Tyr His Ile Lys
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314 Val Ile Asn Ser Ser Ala Arg Arg Ile Gly Tyr Gly Ile Lys Thr Thr
             35 .
315
                                 40
317 Asn Met Lys Arg Leu Gly Val Asp Pro Pro Cys Gly Val Leu Asp Pro
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         50
320 Lys Glu Ala Val Leu Leu Ala Val Ser Cys Asp Ala Phe Ala Phe Gly
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321 65
323 Gln Glu Asp Thr Asn Asn Asp Arg Ile Thr Ile Glu Trp Thr Asn Thr
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326 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
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345 Ile Thr Asn Ala Gly Gly Arg Arg Ile Gly Trp Ala Ile Lys Thr Thr
346
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348 Asn Met Arg Arg Leu Ser Val Asp Pro Pro Cys Gly Val Leu Asp Pro
351 Lys Glu Lys Val Leu Met Ala Val Ser Cys Asp Thr Phe Asn Ala Ala
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354 Thr Glu Asp Leu Asn Asn Asp Arg Ile Thr Ile Glu Trp Thr Asn Thr
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355
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357 Pro Asp Gly Ala Ala Lys Gln Phe Arg Arg Glu Trp Phe Gln Gly Asp
358
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360 Gly Met Val Arg Arg Lys Asn Leu Pro Ile Glu Tyr Asn Leu
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/863,063

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